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                   Welcome to STN International
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NEWS 1
                 Web Page for STN Seminar Schedule - N. America
NEWS 2
         JAN 02
                 STN pricing information for 2008 now available
NEWS 3 JAN 16
                 CAS patent coverage enhanced to include exemplified
                 prophetic substances
NEWS 4
         JAN 28
                 USPATFULL, USPAT2, and USPATOLD enhanced with new
                 custom IPC display formats
NEWS 5 JAN 28 MARPAT searching enhanced
NEWS 6 JAN 28 USGENE now provides USPTO sequence data within 3 days
                 of publication
NEWS 7 JAN 28
                 TOXCENTER enhanced with reloaded MEDLINE segment
NEWS 8 JAN 28 MEDLINE and LMEDLINE reloaded with enhancements
NEWS 9 FEB 08 STN Express, Version 8.3, now available
NEWS 10 FEB 20 PCI now available as a replacement to DPCI
NEWS 11 FEB 25 IFIREF reloaded with enhancements
                 IMSPRODUCT reloaded with enhancements
NEWS 12 FEB 25
NEWS 13 FEB 29 WPINDEX/WPIDS/WPIX enhanced with ECLA and current
                 U.S. National Patent Classification
NEWS 14 MAR 31
                 IFICDB, IFIPAT, and IFIUDB enhanced with new custom
                 IPC display formats
NEWS 15 MAR 31 CAS REGISTRY enhanced with additional experimental
NEWS 16 MAR 31
                 CA/CAplus and CASREACT patent number format for U.S.
                 applications updated
NEWS 17 MAR 31 LPCI now available as a replacement to LDPCI
NEWS 18 MAR 31 EMBASE, EMBAL, and LEMBASE reloaded with enhancements
NEWS 19 APR 04 STN AnaVist, Version 1, to be discontinued
NEWS 20 APR 15 WPIDS, WPINDEX, and WPIX enhanced with new
                 predefined hit display formats
NEWS 21 APR 28 EMBASE Controlled Term thesaurus enhanced
NEWS 22 APR 28 IMSRESEARCH reloaded with enhancements
NEWS 23 MAY 30 INPAFAMDB now available on STN for patent family
                 searching
NEWS 24 MAY 30 DGENE, PCTGEN, and USGENE enhanced with new homology
                 sequence search option
NEWS EXPRESS FEBRUARY 08 CURRENT WINDOWS VERSION IS V8.3,
             AND CURRENT DISCOVER FILE IS DATED 20 FEBRUARY 2008
NEWS HOURS
              STN Operating Hours Plus Help Desk Availability
NEWS LOGIN
              Welcome Banner and News Items
              For general information regarding STN implementation of IPC 8
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Enter NEWS followed by the item number or name to see news on that specific topic.

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FILE 'HOME' ENTERED AT 11:58:57 ON 05 JUN 2008

=> index bioscience medicine

FILE 'DRUGMONOG' ACCESS NOT AUTHORIZED

COST IN U.S. DOLLARS SINCE FILE TOTAL ENTRY SESSION

FULL ESTIMATED COST 0.21 0.21

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ... 'ENTERED AT 11:59:20 ON 05 JUN 2008

72 FILES IN THE FILE LIST IN STNINDEX

Enter SET DETAIL ON to see search term postings or to view search error messages that display as 0* with SET DETAIL OFF.

=> s dicer? or (dsrna?(s) (degrad? or cleav?)) or

((doubl?(s)strand?(s)rna?)(s)(degrad? or cleav?)) or (herna?(s)(degrad? or cleav?))

FILE ADISCTI

8 FILE ADISINSIGHT

368 FILE AGRICOLA

1 FILE ANABSTR

Ω FILE ANTE

FILE AQUALINE

86 FILE AQUASCI

236 FILE BIOENG

2334 FILE BIOSIS

1314 FILE BIOTECHABS

1314 FILE BIOTECHDS

660 FILE BIOTECHNO

13 FILES SEARCHED...

FILE CABA 729

> 1956 FILE CAPLUS

30 FILE CEABA-VTB

10 FILE CIN

32 FILE CONFSCI

FILE CROPB

FILE CROPU

FILE DDFB

FILE DDFU

33 69145 FILE DGENE

23 FILES SEARCHED...

212 FILE DISSABS

FILE DRUGB

FILE DRUGU

87

FILE EMBAL 38

FILE EMBASE 1036

1361 FILE ESBIOBASE

FILE FROSTI

FILE FSTA 31

24883 FILE GENBANK

35 FILES SEARCHED...

1 FILE HEALSAFE

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852
           FILE IFIPAT
            FILE IMSPRODUCT
         1
           FILE IMSRESEARCH
      1510
           FILE LIFESCI
           FILE MEDLINE
      1301
        16
           FILE NTIS
           FILE OCEAN
        21
       626
           FILE PASCAL
       113
           FILE PCTGEN
           FILE PHAR
           FILE PHARMAML
        14
            FILE PHIN
       402
            FILE PROMT
 53 FILES SEARCHED...
           FILE RDISCLOSURE
      1729
            FILE SCISEARCH
       281
           FILE TOXCENTER
           FILE USGENE
      5100
     12510
           FILE USPATFULL
       212
            FILE USPATOLD
           FILE USPAT2
      1890
            FILE VETB
            FILE VETU
        15
            FILE WATER
      1171
            FILE WPIDS
            FILE WPIFV
        14
      1171
             FILE WPINDEX
 69 FILES SEARCHED...
        11
            FILE IPA
         6
             FILE NAPRALERT
           FILE NLDB
        97
 61 FILES HAVE ONE OR MORE ANSWERS, 72 FILES SEARCHED IN STNINDEX
    QUE DICER? OR (DSRNA?(S) (DEGRAD? OR CLEAV?)) OR ((DOUBL?(S) STRAND?(S) RNA?
        )(S)(DEGRAD? OR CLEAV?)) OR (HERNA?(S)(DEGRAD? OR CLEAV?))
=> d rank
        69145
              DGENE
        24883 GENBANK
        12510 USPATFULL
        5100 USGENE
        2334
              BIOSIS
        1956
              CAPLUS
        1890 USPAT2
        1729
              SCISEARCH
        1510
              LIFESCI
        1361
              ESBIOBASE
              BIOTECHABS
        1314
         1314
              BIOTECHDS
               MEDLINE
         1301
         1171
               WPIDS
               WPINDEX
        1171
         1036
               EMBASE
         852
               IFIPAT
         729
               CABA
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F2

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BIOTECHNO

PASCAL

PROMT

BIOENG

281 TOXCENTER

AGRICOLA

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F25
   212 DISSABS
F26
       212 USPATOLD
F27
        113 PCTGEN
F28
        97 NLDB
F29
         87 DRUGU
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         86 AQUASCI
F31
         70 FROSTI
         38 EMBAL
F32
F33
         33 DDFU
F34
         32 CONFSCI
F35
         31 FSTA
F36
         30 CEABA-VTB
F37
         21 OCEAN
F38
         16 NTTS
F39
         1.5
            VETU
F40
         14 PHIN
F41
         14
            WPTFV
         11
F42
             IPA
F43
         10
            CIN
F44
         8 ADISINSIGHT
         8 ANTE
F45
         6
F46
            CROPU
         6
            WATER
F47
         6 NAPRALERT
F48
            DDFB
F49
F50
         5 DRUGB
F51
         2 AQUALINE
F52
         2 VETB
F53
         1 ADISCTI
F54
         1 ANABSTR
F55
          1 CROPB
F56
         1 HEALSAFE
F57
          1 IMSPRODUCT
F58
         1 IMSRESEARCH
F59
         1 PHAR
F60
         1 PHARMAML
F61
         1 RDISCLOSURE
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=> file f2-f13 COST IN U.S. DOLLARS

SINCE FILE TOTAL ENTRY SESSION 5.85

6.06

FULL ESTIMATED COST

FILE 'GENBANK' ENTERED AT 12:04:43 ON 05 JUN 2008

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FILE 'BIOTECHDS' ENTERED AT 12:04:43 ON 05 JUN 2008
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FILE 'MEDITNE' ENTERED AT 12:04:43 ON 05 JUN 2008
=> s dicer? or(dsrna?(s)(degrad? or cleav?)) or
((doubl?(s)strand?(s)rna?)(s)(degrad? or cleav?)) or (herna?(s)(degrad? or cleav?))
   9 FILES SEARCHED...
        55888 DICER? OR(DSRNA?(S)(DEGRAD? OR CLEAV?)) OR ((DOUBL?(S) STRAND?(S
               ) RNA?)(S)(DEGRAD? OR CLEAV?)) OR (HERNA?(S)(DEGRAD? OR CLEAV?))
=> s 12(s)(huma? or sapien?)
   7 FILES SEARCHED...
         5818 L2(S) (HUMA? OR SAPIEN?)
=> s 13(s)(rnase#)
          637 L3(S)(RNASE#)
=> s 14(s)(bindin##)
          182 L4(S)(BINDIN##)
L5
=> dup rem 15
DUPLICATE IS NOT AVAILABLE IN 'GENBANK, USGENE'.
ANSWERS FROM THESE FILES WILL BE CONSIDERED UNIQUE
PROCESSING COMPLETED FOR L5
           160 DUP REM L5 (22 DUPLICATES REMOVED)
=> d ti 16 1-160
    ANSWER 1 OF 160 USPATFULL on STN
L6
TΙ
      System for High Production of Natural and Personalized Interferons
     ANSWER 2 OF 160 USPATFULL on STN
1.6
ΤI
      OLD-35 as an inflammatory agent
     ANSWER 3 OF 160 USPATFULL on STN
L6
      Methods and Materials Relating to Breast Cancer Diagnosis
                                COPYRIGHT 2008 CSA on STN
L6
    ANSWER 4 OF 160 LIFESCI
    Mutations in the U5 Region Adjacent to the Primer Binding Site Affect tRNA
     Cleavage by Human Immunodeficiency Virus Type 1 Reverse Transcriptase In
     Vivo
    ANSWER 5 OF 160 USPATFULL on STN
ΤТ
     Human RNase H1 oligonucleotide compositions thereof
    ANSWER 6 OF 160 USPATFULL on STN
1.6
TΙ
     Methods and compositions for the specific inhibition of gene expression
```

by double-stranded RNA

- L6 ANSWER 7 OF 160 USPATFULL on STN
- II Methods to treat or prevent hormone-resistant prostate cancer using siRNA specific for protocadherin-PC, or other inhibitors of protocadherin-PC expression or activity
- L6 ANSWER 8 OF 160 USPATFULL on STN
- TI Identification of aging genes through large-scale analysis
- L6 ANSWER 9 OF 160 USPATFULL on STN
- TI Compositions and methods for generating short double-stranded rna using mutated rnase III
- L6 ANSWER 10 OF 160 USPATFULL on STN
- TI Primers for synthesizing full-length cDNA and their use
- L6 ANSWER 11 OF 160 USPATFULL on STN
- TI Methods of degrading dsrna and synthesizing rna
- L6 ANSWER 12 OF 160 USPATFULL on STN
- II RNA interference mediating small RNA molecules
- L6 ANSWER 13 OF 160 USPATFULL on STN
- TI BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VACCINIA REGULATORY GENES AND USES THEREOF
- L6 ANSWER 14 OF 160 USPATFULL on STN
- TI Preparation of antibody or an antibody fragment-targeted immunoliposomes for systemic administration of therapeutic or diagnostic agents and uses thereof
- L6 ANSWER 15 OF 160 USPATFULL on STN
- TI Anti-pathogen treatments
- L6 ANSWER 16 OF 160 USPATFULL on STN
- TI BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VACCINIA REGULATORY GENES AND USES THEREOF
- L6 ANSWER 17 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI New capture probe comprising a first adapter segment, a second adapter segment and an miRNA binding segment, useful for isolating miRNAs;

 DNA probe capture and immobilization for microRNA isolation
- L6 ANSWER 18 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 1
- TI Small-molecule activators of RNase L with broad-spectrum antiviral activity
- L6 ANSWER 19 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 2
- TI Identification of the gene encoding a type 1 RNase H with an N-terminal double-stranded RNA binding domain from a psychrotrophic bacterium
- L6 ANSWER 20 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 3
- TI Proteomic and functional analysis of Argonaute-containing mRNA-protein complexes in human cells
- L6 ANSWER 21 OF 160 Elsevier BIOBASE COPYRIGHT 2008 Elsevier Science B.V.
- TI Identification and biochemical analysis of a mitochondrial endonuclease of Podospora anserina related to curved-DNA binding proteins
- L6 ANSWER 22 OF 160 CAPLUS COPYRIGHT 2008 ACS on STN

- TI Evolutionary conservation of a unique amino acid sequence in human DICER protein essential for binding to Argonaute family proteins
- L6 ANSWER 23 OF 160 Elsevier BIOBASE COPYRIGHT 2008 Elsevier Science B.V. on STN
- TI Homodimeric Structure and Double-stranded RNA Cleavage Activity of the C-terminal RNase III Domain of Human Dicer
- L6 ANSWER 24 OF 160 USPATFULL on STN
- TI Methods for the identification, assessment, and treatment of patients with cancer therapy
- L6 ANSWER 25 OF 160 USPATFULL on STN
- TI Oligomeric compounds and compositions for use in modulation of pri-mirnas
- L6 ANSWER 26 OF 160 USPATFULL on STN
- TI Identification of gene expression by heart failure etiology
- L6 ANSWER 27 OF 160 USPATFULL on STN
- TI Cloning and characterization of microRNAs from rice
- L6 ANSWER 28 OF 160 USPATFULL on STN
- TI Dicer interacting proteins and uses therefor
- L6 ANSWER 29 OF 160 USPATFULL on STN
- TI Soluble rna polymerase protein and methods for the use thereof
- L6 ANSWER 30 OF 160 USPATFULL on STN
- TI Allele-specific RNA interference
- L6 ANSWER 31 OF 160 USPATFULL on STN
- TI Pharmaceutical compositions and methods useful for modulating angiogenesis, inhibiting metastasis and tumor fibrosis, and assessing the malignancy of colon cancer tumors
- L6 ANSWER 32 OF 160 USPATFULL on STN
- TI Genes displaying enhanced expression during cellular senescence and terminal cell differentiation and uses thereof
- L6 ANSWER 33 OF 160 USPATFULL on STN
- TI Methods and compositions for generating recombinant nucleic acid molecules
- L6 ANSWER 34 OF 160 USPATFULL on STN
- TI Means and methods for the specific modulation of target genes in the cns and the eye and methods for their identification
- L6 ANSWER 35 OF 160 USPAT2 on STN
- TI DICER INTERACTING PROTEINS AND USES THEREFOR
- L6 ANSWER 36 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Composition for inhibiting proliferation of lung cancer exhibiting neuroendocrine differentiation, contains vector having DNA encoding small interfering RNA having sequence of human achaete scute homologue 1 mRNA, and carrier:

vector-mediated small interfering RNA-encoding DNA transfer and expression in neuroendocrine differentiation-positive lung cancer for neuroendocrine differentiation-positive lung cancer proliferation inhibition and gene therapy

- TI DUF283 domain of Dicer proteins has a double-stranded RNA-binding fold
- L6 ANSWER 38 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 5
- TI The role of PACT in the RNA silencing pathway
- L6 ANSWER 39 OF 160 USPATFULL on STN
- TI Polynucleotide compositions encoding S-adenosyl-L
 - methionine:phosphoethanolamine N-methyltransferase and methods for modulating lipid biosynthesis in plants
- L6 ANSWER 40 OF 160 USPATFULL on STN
- TI Modular design and construction of nucleic acid molecules,
 - aptamer-derived nucleic acid constructs, RNA scaffolds, their expression, and methods of use
- L6 ANSWER 41 OF 160 USPATFULL on STN
- TI Oligomeric compounds and compositions for use in modulation small non-coding RNAs
- L6 ANSWER 42 OF 160 USPATFULL on STN
- TI Oligo-or polynucleotides
- L6 ANSWER 43 OF 160 USPATFULL on STN
- TI Composition and method for introduction of RNA interference sequences into targeted cells and tissues
- L6 ANSWER 44 OF 160 USPATFULL on STN
- TI RNA interference mediating small RNA molecules
- L6 ANSWER 45 OF 160 USPATFULL on STN
- TI Means and methods for the specific inhibition of genes in cells and tissue of the cns and/or eye
- L6 ANSWER 46 OF 160 USPATFULL on STN
- TI Methods and compositions for enhancing the efficacy and specificity of RNAi
- L6 ANSWER 47 OF 160 USPATFULL on STN
- TI Methods and compositions for enhancing the efficacy and specificity of RNAi
- L6 ANSWER 48 OF 160 USPATFULL on STN
- TI Methods of using mammalian RNase H and compositions thereof
- L6 ANSWER 49 OF 160 USPATFULL on STN
- TI Human RNase III and compositions and uses thereof
- L6 ANSWER 50 OF 160 USPATFULL on STN
- TI Cell death-related nucleases and their uses
- L6 ANSWER 51 OF 160 USPATFULL on STN
- TI Compositions and methods for preparing short RNA molecules and other nucleic acids
- L6 ANSWER 52 OF 160 USPATFULL on STN
- TI Acyl-nucleotide probes and methods of their synthesis and use in proteomic analysis
- L6 ANSWER 53 OF 160 USPATFULL on STN
- TI Syndecans and angiogenesis
- L6 ANSWER 54 OF 160 USPATFULL on STN

- TI DECREASING GENE EXPRESSION IN A MAMMALIAN SUBJECT IN VIVO VIA AAV-MEDIATED RNA1 EXPRESSION CASSETTE TRANSFER
- L6 ANSWER 55 OF 160 USPAT2 on STN

TI

- Cell death-related nucleases and their uses
- L6 ANSWER 56 OF 160 USPAT2 on STN
- TI Acyl-nucleotide probes and methods of their synthesis and use in proteomic analysis
- L6 ANSWER 57 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
 TI New composition comprising modified nuclease, useful for treating or
 preventing a viral disease or a non-viral disease, e.g. Alzheimer
 disease, Parkinson disease, multiple sclerosis or age-related dementia;
 the use of a recombinant nuclease and peptide nucleic acid in a
 composition for a neurodegenerative disease gene therapy application
- L6 ANSWER 58 OF 160 BIOTECHOS COPYRIGHT 2008 THOMSON REUTERS on STN

 New oligomeric compound that can hybridize with or sterically interfere
 with nucleic acid molecules comprising or encoding small non-coding RNA
 targets, useful for treating e.g., cancer and diabetes;
 vector-mediated human ERK5 protein-specific small interfering RNA and
 antisense oligonucleotide administration and expression in stem cell
 for use in disease gene therapy and RNA interference
- L6 ANSWER 59 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Inactivating a virus (e.g. HCV) in a patient to treat the viral infection comprises administering to the patient a modified small interfering RNA in an amount to inactivate the virus; virus replication inhibition using RNA interference for use in gene

therapy

- L6 ANSWER 60 OF 160 Elsevier BIOBASE COPYRIGHT 2008 Elsevier Science B.V. on STN
- TI Normal microRNA maturation and germ-line stem cell maintenance requires loquacious, a double-stranded RNA-binding domain protein
- L6 ANSWER 61 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 6
- $\ensuremath{\mathsf{TI}}$ TRBP recruits the Dicer complex to Ago2 for microRNA processing and gene silencing
- L6 ANSWER 62 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 7
- TI Structural basis for 5'-end-specific recognition of guide RNA by the A. fulgidus Piwi protein
- L6 ANSWER 63 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN
- TI Normal microRNA Maturation and Germ-Line Stem Cell Maintenance Requires Loquacious, a Double-Stranded RNA-Binding Domain Protein
- L6 ANSWER 64 OF 160 USPATFULL on STN
- TI Methods of using mammalian RNase H and compositions thereof
- L6 ANSWER 65 OF 160 USPATFULL on STN
- TI Modulation of PAI-1 mRNA-binding protein expression
- L6 ANSWER 66 OF 160 USPATFULL on STN
- TI Expression profiles for breast cancer and methods of use
- L6 ANSWER 67 OF 160 USPATFULL on STN
- TI Oligoribonucleotides and ribonucleases for cleaving RNA
- L6 ANSWER 68 OF 160 USPATFULL on STN

- TI Human RNase III and compositions and uses thereof
- L6 ANSWER 69 OF 160 USPATFULL on STN
- TI Human RNase III and compositions and uses thereof
- L6 ANSWER 70 OF 160 USPATFULL on STN
- TI Human RNase H1 and oligonucleotide compositions thereof
- L6 ANSWER 71 OF 160 USPATFULL on STN
- TI Anti-pathogen treatments
- L6 ANSWER 72 OF 160 USPAT2 on STN
- TI RNA interference mediating small RNA molecules
- L6 ANSWER 73 OF 160 USPAT2 on STN
 - RNA interference mediating small RNA molecules
- L6 ANSWER 74 OF 160 USPAT2 on STN
- TI Anti-pathogen treatments

ΤI

- L6 ANSWER 75 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI New RNase resistant small interfering RNA, useful for treating viral infections, e.g., hepatitis C, influenza virus or coronavirus infection; small interfering RNA transfer and expression in host cell for RNA interference and gene therapy
- L6 ANSWER 76 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Making a cell or multicellular organism with resistance to a viral pathogen susceptible to infect the cell or organism by genetically modifying the cell or organism to utilize polynucleic acid molecule or viral RNA transcript as a template;
 - transgenic plant construction using genetically modified cell and polynucleic acid molecule for use in disease-resistance
- L6 ANSWER 77 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 8
- TI Two Modes of HIV-1 Polypurine Tract Cleavage Are Affected by Introducing Locked Nucleic Acid Analogs into the (-) DNA Template
- L6 ANSWER 78 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 9
- TI The Drosha-DGCR8 complex in primary microRNA processing
- L6 ANSWER 79 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 10
- TI Single Processing Center Models for Human Dicer and Bacterial RNase III
- L6 ANSWER 80 OF 160 USPATFULL on STN
- TI Human Rnase H1 and oligonucleotide compositions thereof
- L6 ANSWER 81 OF 160 USPATFULL on STN
- TI MDA-7 proteins and methods of use thereof
- L6 ANSWER 82 OF 160 USPATFULL on STN
- TI Syndecans and angiogenesis
- L6 ANSWER 83 OF 160 USPATFULL on STN
- TI 207 human secreted proteins
- L6 ANSWER 84 OF 160 USPATFULL on STN
- TI Compositions and methods for the therapy and diagnosis of colon cancer
- L6 ANSWER 85 OF 160 USPATFULL on STN
- TI Oligoribonucleotides and ribonucleases for cleaving RNA

- L6 ANSWER 86 OF 160 USPATFULL on STN
- TI Genes displaying enhanced expression during cellular senescence and terminal cell differentiation and uses thereof
- L6 ANSWER 87 OF 160 USPATFULL on STN
- TI Oligoribonucleotides and ribonucleases for cleaving RNA
- L6 ANSWER 88 OF 160 USPATFULL on STN
- TI Oligoribonucleotides and ribonucleases for cleaving RNA
- L6 ANSWER 89 OF 160 USPATFULL on STN
- TI Oligoribonucleotides and ribonucleases for cleaving RNA
- L6 ANSWER 90 OF 160 USPATFULL on STN
- TI Compositions and methods for the therapy and diagnosis of pancreatic cancer
- L6 ANSWER 91 OF 160 USPATFULL on STN
- TI Human RNase III and compositions and uses thereof
- L6 ANSWER 92 OF 160 USPAT2 on STN
- TI Genes displaying enhanced expression during cellular senescence and terminal cell differentiation and uses thereof
- L6 ANSWER 93 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Reducing expression of target gene in cell, by incubating dsRNA corresponding to part of target gene with effective amount of composition comprising RNase III domain, and transfecting siRNA into cell; involving DNA primer, polymerase chain reaction and RNA-polymerase
- L6 ANSWER 94 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI New nucleobase oligomers that inhibit expression of inhibitor of apoptosis gene, useful for treating cancer and other lymphoproliferative disorders by inducing apoptosis;
 - antisense oligonucleotide or dsRNA transfer and expression in host cell for cancer gene therapy
- L6 ANSWER 95 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 11
- TI Degradation of Double-Stranded RNA by Human Pancreatic Ribonuclease: Crucial Role of Noncatalytic Basic Amino Acid Residues
- L6 ANSWER 96 OF 160 CAPLUS COPYRIGHT 2008 ACS on STN
- TI Identification of eight members of the Argonaute family in the human genome
- L6 ANSWER 97 OF 160 USPATFULL on STN
- TI METHOD FOR GENERATING A SUBTRACTED CDNA LIBRARY AND USES OF THE GENERATED LIBRARY
- L6 ANSWER 98 OF 160 USPATFULL on STN
- TI Oligoribonucleotides and ribonucleases for cleaving RNA
- L6 ANSWER 99 OF 160 USPATFULL on STN
- II Human RNase III and compositions and uses thereof
- L6 ANSWER 100 OF 160 USPATFULL on STN
- TI Human RNase H1 mutants
- L6 ANSWER 101 OF 160 USPATFULL on STN
- TI Compositions and methods for the therapy and diagnosis of colon cancer
- L6 ANSWER 102 OF 160 USPATFULL on STN

- TI Compositions and methods for the therapy and diagnosis of ovarian cancer
- L6 ANSWER 103 OF 160 USPAT2 on STN
- TI MDA-7 nucleic acid molecules and pharmaceutical compositions thereof
- L6 ANSWER 104 OF 160 USPAT2 on STN
- TI Human RNase III and compositions and uses thereof
- L6 ANSWER 105 OF 160 USPAT2 on STN
- TI Human RNase H1 mutants
- L6 ANSWER 106 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Novel dynamic action reference tool (DART) comprising a molecular shaft covalently linked to a linkage polypeptide covalently linked to a molecular point, useful for isolating and analyzing nucleic acids, polypeptides;
 - dynamic action reference tool and vector expression in host cell for use in disease gene therapy
- L6 ANSWER 107 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
 TI Novel human RNase H1 polypeptide comprising mutations compared to wild
- type human RNase HI useful for inhibiting expression of selected protein by antisense oligonucleotide targeted to RNA encoding selected protein; vector plasmid pETI7b-mediated gene transfer and expression in host cell for use in gene therapy
- L6 ANSWER 108 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Generating mRNA-CDNA hybrids for suppressing cancer-related genes, or treating or preventing microbe related genes, comprises thermocycling steps of promoter-linked double-stranded cDNA or RNA synthesis;
 - useful for gene therapy, high throughput screening, DNA microarray analysis and functional genomics
- L6 ANSWER 109 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN Regulating gene expression in plants for controlling gene silencing, comprises altering the transcription or translation of an endonuclease nucleotide sequence encoding a polypeptide comprising an exonuclease
 - domain; Arabidopsis sp. transgenic plant and seed construction involving vector-mediated beta-glucuronidase gene transfer and expression in plant cell
- L6 ANSWER 110 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Probes for detecting target nucleotide sequence in sample, has sequence that forms hairpin structure having a double-stranded segment and single-stranded loop collectively forming region complementary to target sequence;
 - oligonuclectide DNA probe, RNA probe, peptide nucleic acid probe for detecting target sequence in a sample and for transcription and/or DNA amplification of probe sequence
- L6 ANSWER 111 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI New synthetic oligomeric compound that is specifically hybridizable with a preselected RNA target, useful for treating an organism having a disease characterized by the undesired production of a protein;
 - RNA-specific oligonucleotide transfer and expression in host cell for gene therapy
- L6 ANSWER 112 OF 160 Elsevier BIOBASE COPYRIGHT 2008 Elsevier Science B.V. on STN
- TI Human Dicer preferentially cleaves dsRNAs at their termini without a requirement for ATP

- L6 ANSWER 113 OF 160 CAPLUS COPYRIGHT 2008 ACS on STN
- TI Ribonuclease activity and RNA binding of recombinant human Dicer
- L6 ANSWER 114 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 12
- TI Control of developmental timing by small temporal RNAs: a paradigm for RNA-mediated regulation of gene expression
- L6 ANSWER 115 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN
- TI Mouse ribonuclease III. cDNA structure, expression analysis, and chromosomal location
- L6 ANSWER 116 OF 160 USPATFULL on STN
- TI Virus resistant plants expressing animal cell-derived
 - (2'-5')oligadenylate synthetase and ribonuclease L and A method for creating the same
- L6 ANSWER 117 OF 160 BIOTECHDS COPYRIGHT 2008 THOMSON REUTERS on STN
- TI Cobra Venom Factor 1 polypeptides which are analogs of Complement Component C3, useful for identifying compounds that may be used to modulate the complement system;
 - snake venom production involving vector plasmid pSPORT-mediated gene transfer for expression in host cell e.g. Escherichia coli
- L6 ANSWER 118 OF 160 BIOSIS COPYRIGHT (c) 2008 The Thomson Corporation on STN
- TI Investigating the structure of human RNase H1 by site-directed mutagenesis.
- L6 ANSWER 119 OF 160 USPATFULL on STN
- TI Oligoribonucleotides and ribonucleases for cleaving RNA
- L6 ANSWER 120 OF 160 USPATFULL on STN TI Uses of mda-6

ΤТ

- L6 ANSWER 121 OF 160 USPATFULL on STN
- TI Animal 2-5A-dependent RNases and encoding sequences therefor
- L6 ANSWER 122 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 14
- TI Activation of the Interferon-Inducible (2'-5') Oligoadenylate Synthetase by the Epstein-Barr Virus RNA, EBER-1
- L6 ANSWER 123 OF 160 USPATFULL on STN
- TI Human 26S proteasome subunit components
- L6 ANSWER 124 OF 160 USPATFULL on STN
 - Peptidyl derivatives as inhibitors of pro-apoptotic cysteine proteinases
- L6 ANSWER 125 OF 160 Elsevier BIOBASE COPYRIGHT 2008 Elsevier Science B.V. on STN
- TI The 2-5A system: Modulation of viral and cellular processes through acceleration of RNA degradation
- L6 ANSWER 126 OF 160 USPATFULL on STN
- $\ensuremath{\mathsf{TI}}$ Method for generating a subtracted cDNA library and uses of the generated library
- L6 ANSWER 127 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 15
- TI Catalytic cleavage of an RNA target by 2-5A antisense and RNase L
- L6 ANSWER 128 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN
- TI Induction of apoptotic nuclei by interferon- gamma and by predesquamin in

cultured keratinocytes

cultured keratinocytes
L6 ANSWER 129 OF 160 SCISEARCH COPYRIGHT (c) 2008 The Thomson Corporation on STN DUPLICATE 16
TI HIV-1 REVERSE TRANSCRIPTASE-ASSOCIATED RNASE-H CLEAVES RNA/RNA IN ARRESTED COMPLEXES - IMPLICATIONS FOR THE MECHANISM BY WHICH RNASE-H DISCRIMINATES BEWEEN RNA/RNA AND RNA/DNA
L6 ANSWER 130 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN TI Localization of the interferon-induced, 2-5A-Dependent RNase gene (RNS4) to human chromosome 1q25
L6 ANSWER 131 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN TI Double-stranded RNA-dependent RNase activity associated with human immundeficiency virus type 1 reverse transcriptase.
L6 ANSWER 132 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): Genomic Islands in the Pathogenic Filamentous Fungus Aspergillus fumigatus TITLE (TI): Direct Submission
iiile (ii): Direct Submission
L6 ANSWER 133 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): Genomic analysis of Bartonella identifies type IV secretion systems as host adaptability factors
TITLE (II): Direct Submission
L6 ANSWER 134 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): Complete sequence of Clostridium phytofermentans ISDg TITLE (TI): Direct Submission
L6 ANSWER 135 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): The genome sequence of Schizosaccharomyces pombe TITLE (TI): Direct Submission
L6 ANSWER 136 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): The Chlamydomonas genome reveals the evolution of key animal and plant functions
TITLE (TI): Direct Submission
L6 ANSWER 137 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): The genome sequence of Schizosaccharomyces pombe TITLE (TI): Direct Submission
L6 ANSWER 138 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): Comparative genomics of trypanosomatid parasitic protozoa
TITLE (TI): The Genome of the African Trypanosome Trypanosoma brucei
TITLE (TI): Direct Submission
L6 ANSWER 139 OF 160 GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): Genome sequence of a proteolytic (Group I) Clostridium botulinum strain Hall A and comparative analysis of the clostridial genomes

clostridial genomes

TITLE (TI):	Direct Submission
L6 ANSWER 140 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI):	Comparative genomics of trypanosomatid parasitic protozoa
TITLE (TI):	The Genome of the African Trypanosome Trypanosoma brucei
TITLE (TI):	Sequencing, closure, and annotation of Trypanosoma brucei chromosomes 2 through 8
TITLE (TI):	Direct Submission
L6 ANSWER 141 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI):	Comparative genomics of trypanosomatid parasitic protozoa
TITLE (TI):	The Genome of the African Trypanosome Trypanosoma
TITLE (TI):	Sequencing, closure, and annotation of Trypanosoma brucei chromosomes 2 through 8
TITLE (TI):	Direct Submission
L6 ANSWER 142 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI):	Annotation and evolutionary relationships of a small regulatory RNA gene micF and its target ompF in
TITLE (TI):	Yersinia species The Complete Genome Sequence and Comparative Genome Analysis of the High Pathogenicity Yersinia
TITLE (TI):	enterocolitica Strain 8081 Direct Submission
L6 ANSWER 143 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI):	Complete DNA sequence of a serogroup A strain of Neisseria meningitidis Z2491
TITLE (TI):	Direct Submission
L6 ANSWER 144 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI):	Complete genome of the mutualistic, N2-fixing grass endophyte Azoarcus sp. strain BH72
TITLE (TI):	Direct Submission
L6 ANSWER 145 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI):	The multidrug-resistant human pathogen Clostridium difficile has a highly mobile, mosaic genome
TITLE (TI):	The multidrug resistant human pathogen Clostridium difficile has a highly mobile, mosaic genome
TITLE (TI):	Direct Submission
L6 ANSWER 146 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI):	Complete Sequence of Chromosome 1 of Rhodobacter sphaeroides 2.4.1
TITLE (TI):	Sphaeroides 2.4.1 Direct Submission
L6 ANSWER 147 OF 160	GENBANK® COPYRIGHT 2008 on STN
TITLE (TI): TITLE (TI):	Complete sequence of Psychrobacter arcticum 273-4 Direct Submission

L6	ANSWER 148	OF	160	GENBANK® COPYRIGHT 2008 on STN
	TITLE (TI):			Extensive DNA inversions in the B. fraqilis genome
	TITLE (TI):			control variable gene expression Direct Submission
1.0		OF	160	
L6	ANSWER 149	OF	100	
	TITLE (TI): TITLE (TI):			The map-based sequence of the rice genome The Rice Annotation Project Database (RAP-DB): hub for Oryza sativa ssp. japonica genome information
	TITLE (TI):			Curated genome annotation of Oryza sativa ssp. japonica and comparative genome analysis with Arabidopsis thaliana
	TITLE (TI):			Oryza sativa nipponbare(GA3) genomic DNA, chromosome 3
	TITLE (TI): TITLE (TI):			The First Rice Annotation Project Meeting (RAP1) Direct Submission
L6	ANSWER 150	OF	160	GENBANK® COPYRIGHT 2008 on STN
	TITLE (TI):			The genome of the heartwater agent Ehrlichia ruminantium contains multiple tandem repeats of
	TITLE (TI):			actively variable copy number Direct Submission
L6	ANSWER 151	OF	160	GENBANK® COPYRIGHT 2008 on STN
	TITLE (TI):			Genome sequence of Yersinia pestis, the causative agent of plaque
	TITLE (TI):			Annotation and evolutionary relationships of a small regulatory RNA gene micF and its target ompF in Yersinia species
	TITLE (TI):			Direct Submission
L6	ANSWER 152	OF	160	GENBANK® COPYRIGHT 2008 on STN
	TITLE (TI):			The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hypervariable sequences
	TITLE (TI):			Re-annotation and re-analysis of the Campylobacter jejuni NCTC11168 genome sequence
	TITLE (TI): TITLE (TI):			Direct Submission Direct Submission
L6	ANSWER 153	OF	160	GENBANK® COPYRIGHT 2008 on STN
	TITLE (TI):			Genomic plasticity of the causative agent of melioidosis, Burkholderia pseudomallei
	TITLE (TI):			Direct Submission
L6	ANSWER 154	OF	160	GENBANK® COPYRIGHT 2008 on STN
	TITLE (TI): TITLE (TI):			Genome evolution in yeasts Direct Submission
L6	ANSWER 155	OF	160	GENBANK® COPYRIGHT 2008 on STN
	TITLE (TI): TITLE (TI):			Genome evolution in yeasts Direct Submission
L6	ANSWER 156	OF	160	GENBANK® COPYRIGHT 2008 on STN

TITLE (TI): Genome evolution in yeasts

TITLE (TI): Direct Submission

L6 ANSWER 157 OF 160 GENBANK® COPYRIGHT 2008 on STN

TITLE (TI): Genome evolution in yeasts

TITLE (TI): Direct Submission

L6 ANSWER 158 OF 160 GENBANK® COPYRIGHT 2008 on STN

TITLE (TI): Genome evolution in yeasts

TITLE (TI): Direct Submission

L6 ANSWER 159 OF 160 GENBANK® COPYRIGHT 2008 on STN

TITLE (TI): Complete genomes of two clinical Staphylococcus aureus

strains: evidence for the rapid evolution of virulence and drug resistance

TITLE (TI): Direct Submission

L6 ANSWER 160 OF 160 GENBANK® COPYRIGHT 2008 on STN

TITLE (TI): Sequencing and analysis of the genome of the Whipple's

disease bacterium Tropheryma whipplei

TITLE (TI): Direct Submission

=> d ibib abs 16 6 9 11 12 22 23 28 37 95 99 113

L6 ANSWER 6 OF 160 USPATFULL on STN

ACCESSION NUMBER: 2007:303223 USPATFULL

TITLE: Methods and compositions for the specific inhibition of

gene expression by double-stranded RNA
INVENTOR(S): Rossi, John J., Alta Loma, CA, UNITED STATES

Behlke, Mark A., Coralville, IA, UNITED STATES

Kim, Dongho, Los Angeles, CA, UNITED STATES
PATENT ASSIGNEE(S): City of Hope, Duarte, CA, UNITED STATES (U.S.

corporation)

Integrated DNA Technologies, Inc., Coralville, IA,

UNITED STATES (U.S. corporation)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 2005-79906, filed

on 15 Mar 2005, PENDING

NUMBER DATE

PRIORITY INFORMATION: US 2004-553487P 20040315 (60)
DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: ROTHWELL, FIGG, ERNST & MANBECK, P.C., 1425 K STREET,

N.W., SUITE 800, WASHINGTON, DC, 20005, US

NUMBER OF CLAIMS: 14 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 31 Drawing Page(s)

LINE COUNT: 4653

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The invention is directed to compositions and methods for selectively reducing the expression of a gene product from a desired target gene in

a cell, as well as for treating diseases caused by the expression of the gene. More particularly, the invention is directed to compositions that contain double stranded RNA ("dsRNA"), and methods for preparing them, that are capable of reducing the expression of target genes in eukaryotic cells. The dsRNA has a first oligonucleotide sequence that is between 25 and about 30 nucleotides in length and a second oligonucleotide sequence that anneals to the first sequence under biological conditions. In addition, a region of one of the sequences of the dsRNA having a sequence length of at least 19 nucleotides is sufficiently complementary to a nucleotide sequence of the RNA produced from the target gene to trigger the destruction of the target RNA by the RNAi machinery.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L6 ANSWER 9 OF 160 USPATFULL on STN

ACCESSION NUMBER: 2007:177863 USPATFULL

TITLE: Compositions and methods for generating short

double-stranded rna using mutated rnase III INVENTOR(S): Maina, Claude V., West Newbury, NY, UNITED STATES Tzertzinis, George, Cambridge, MA, UNITED STATES

Kumar, Sanjay, Ipswich, MA, UNITED STATES New England Biolabs, Inc., Ipswich, MA, UNITED STATES, PATENT ASSIGNEE(S):

01938 (U.S. corporation)

NUMBER KIND DATE US 20070155684 A1 20070705 US 2005-586720 A1 20050121 (10) PATENT INFORMATION: APPLICATION INFO.: WO 2005-US2029 20050121 20060720 PCT 371 date

NUMBER DATE US 2004-538805P 20040123 (60) PRIORITY INFORMATION: US 2004-543880P 20040212 (60)

US 2004-572240P 20040518 (60) DOCUMENT TYPE: Utility

FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: HARRIET M. STRIMPEL, NEW ENGLAND BIOLABS, INC., 240

COUNTY ROAD, IPSWICH, MA, 01938-2723, US

30 NUMBER OF CLAIMS:

EXEMPLARY CLAIM: NUMBER OF DRAWINGS: 17 Drawing Page(s)

LINE COUNT: 1831

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Compositions and methods are provided for preparing an hsiRNA mixture and for silencing of gene expression in vivo. The composition relates to a mutant RnaseIII. The methods are directed to reacting a preparation of dsRNA with an effective amount of a mutant RNAse III to produce the hsiRNA mixture.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L6 ANSWER 11 OF 160 USPATFULL on STN

ACCESSION NUMBER: 2007:120911 USPATFULL

TITLE: Methods of degrading dsrna and synthesizing rna

INVENTOR(S): Sagawa, Hiroaki, Shiga, JAPAN Tomono, Jun, Okayama, JAPAN

Ueno, Harumi, Shiga, JAPAN Kato, Ikunoshin, Shiga, JAPAN

PATENT ASSIGNEE(S): TAKARA BIO INC., Otsu-shi, JAPAN, 520-2193 (non-U.S.

corporation)

	NUMBER	KIND	DATE		
PATENT INFORMATION:	US 20070105113	A1	20070510		
APPLICATION INFO.:	US 2004-567731	A1	20040810	(10)	
	WO 2004-JP11480		20040810		
			20060210	PCT 371	date

			NUMBER	DATE
PRIORITY	INFORMATION:	JP	2003-293553	20030814
		JP	2003-342126	20030930
		JP	2003-409639	20031208
		JP	2004-86129	20040324
DOCUMENT	TYPE:	II+ ·	ility	

DOCUMENT TYPE: Utility

FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: BROWDY AND NEIMARK, P.L.L.C., 624 NINTH STREET, NW,

SUITE 300, WASHINGTON, DC, 20001-5303, US
NUMBER OF CLAIMS: 29

EXEMPLARY CLAIM: 1
LINE COUNT: 3027

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

A protein having an activity of degrading a dsRNA, namely, being capable of acting on a long-chain dsRNA to form a dsRNA of a definite length; a method of efficiently preparing a dsRNA of a definite length which comprises treating a dsRNA with the protein having an activity of degrading a dsRNA in the coexistence of a protein having an activity of binding to a nucleic acid such as a protein having an RNA-binding activity; and a method of using the protein having an activity of binding to a nucleic acid to elevate the efficiency in an RNA synthesis reaction typified by dsRNA synthesis.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L6 ANSWER 12 OF 160 USPATFULL on STN

ACCESSION NUMBER: 2007:107472 USPATFULL

TITLE: RNA interference mediating small RNA molecules

INVENTOR(S): Tuschl, Thomas, New York, NY, UNITED STATES

Elbashir, Sayda Mahgoub, Cambridge, MA, UNITED STATES Lendeckel, Winfried, Hohengandern, GERMANY, FEDERAL

REPUBLIC OF

PATENT ASSIGNEE(S): Max-Planck-Gesellschaft zur Forderung der

Wissenschaften e. V., Munchen, GERMANY, FEDERAL

REPUBLIC OF (non-U.S. corporation)

			NUMBER	KIND	DATE
PATENT	INFORMATION:	US	20070093445	A1	20070426

APPLICATION INFO.: US 2006-634129 A1 20061206 (11)
RELATED APPLN. INFO.: Division of Ser. No. US 2004-433050, filed on 26 Jul

2004, PENDING A 371 of International Ser. No. WO

2001-EP13968, filed on 29 Nov 2001

		NUMBER	DATE	
PRIORITY	INFORMATION:		20001201 20010330 (60)	
DOCUMENT	TYPE:	Utility		

FILE SEGMENT: APPLICATION
LEGAL REPRESENTATIVE: ROTHWELL, FIGG, ERNST & MANBECK, P.C., 1425 K STREET,
N.W., SUITE 800, WASHINGTON, DC, 20005, US

NUMBER OF CLAIMS: 2.9 EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 23 Drawing Page(s)

LINE COUNT:

2309 CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Double-stranded RNA (dsRNA) induces sequence-specific

post-transcriptional gene silencing in many organisms by a process known as RNA interference (RNAi). Using a Drosophila in vitro system, we demonstrate that 19-23 nt short RNA fragments are the sequence-specific mediators of RNAi. The short interfering RNAs (siRNAs) are generated by an RNase III-like processing reaction from long dsRNA. Chemically synthesized siRNA duplexes with overhanging 3' ends mediate efficient target RNA cleavage in the lysate, and the cleavage site is located near the center of the region spanned by the guiding siRNA. Furthermore, we provide evidence that the direction of dsRNA processing determines whether sense or antisense target RNA can be cleaved by the produced siRNP complex.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 22 OF 160 CAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2007:683471 CAPLUS

DOCUMENT NUMBER: 147:252868

TITLE: Evolutionary conservation of a unique amino acid

sequence in human DICER protein essential for binding

to Argonaute family proteins

Sasaki, Takashi; Shimizu, Nobuyoshi AUTHOR(S):

CORPORATE SOURCE: Department of Molecular Biology, Keio University

School of Medicine, Shinjuku-ku, Tokyo, 160-8582,

Japan

SOURCE: Gene (2007), 396(2), 312-320 CODEN: GENED6; ISSN: 0378-1119

PUBLISHER: Elsevier B.V. DOCUMENT TYPE: Journal LANGUAGE: English

The Argonaute family and DICER proteins are major key proteins involved in the RNA-mediated gene silencing mechanism of various species. In this mechanism, cleavage of mRNAs (mRNA) or suppression of mRNA translation takes place via small RNAs that are uniquely processed by DICER. Previously, human Argonaute family proteins were demonstrated to bind to DICER. This study identified a unique amino acid sequence of 127 amino acids in the RIBOc-A domain of human DICER protein as a "binding site" to Argonaute proteins. Comparative genomics anal. revealed that this unique amino acid sequence is highly conserved in the vertebrates, but not found in the non-vertebrate species. Significant difference in the RIBOc-A domain of DICER protein between vertebrate and non-vertebrate species may help exploring the functional complexity in the RNA-mediated gene

silencing mechanism. REFERENCE COUNT: 50 THERE ARE 50 CITED REFERENCES AVAILABLE FOR THIS RECORD, ALL CITATIONS AVAILABLE IN THE RE FORMAT

L6 ANSWER 23 OF 160 Elsevier BIOBASE COPYRIGHT 2008 Elsevier Science B.V. on STN

ACCESSION NUMBER: 2007297926 ESBIOBASE

TITLE: Homodimeric Structure and Double-stranded RNA Cleavage Activity of the C-terminal RNase III Domain of Human Dicer

AUTHOR: Takeshita D.; Zenno S.; Lee W.C.; Nagata K.; Saigo K.; Tanokura M.

CORPORATE SOURCE: M. Tanokura, Department of Applied Biological

Chemistry, Graduate School of Agricultural and Life Sciences, University of Tokyo, 1-1-1 Yayoi, Bunkyo-ku, Tokyo, 113-8657, Japan.

E-mail: amtanok@mail.ecc.u-tokvo.ac.jp Journal of Molecular Biology, (16 NOV 2007), 374/1

(106-120), 48 reference(s)

CODEN: JMOBAK ISSN: 0022-2836

PUBLISHER ITEM IDENT.: S0022283607011680 DOCUMENT TYPE: Journal; Article United Kingdom

LANGUAGE: English SUMMARY LANGUAGE: Enalish

SOURCE:

Human Dicer contains two RNase III domains

(RNase IIIa and RNase IIIb) that are responsible for

the production of short interfering RNAs and microRNAs. These small RNAs induce gene silencing known as RNA

interference. Here, we report the crystal structure of the C-terminal

RNase III domain (RNase IIIb) of human

Dicer at 2.0 Å resolution. The structure revealed that the

RNase IIIb domain can form a tightly associated homodimer, which is similar to the dimers of the bacterial RNase III domains and the two RNase III domains of Giardia Dicer.

Biochemical analysis showed that the RNase IIIb homodimer can

cleave double-stranded RNAs (

dsRNAs), and generate short dsRNAs with 2 nt 3'

overhang, which is characteristic of RNase III products. The RNase IIIb domain contained two magnesium ions per monomer around

the active site. The distance between two Mg-1 ions is approximately 20.6 Å, almost identical with those observed in bacterial RNase III enzymes and Giardia Dicer, while the locations of two Mg-2 ions were not conserved at all. We presume that Mg-1 ions act as catalysts for dsRNA cleavage, while Mg-2 ions are

involved in RNA binding. . COPYRGT. 2007 Elsevier Ltd.

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ANSWER 28 OF 160 USPATFULL on STN

ACCESSION NUMBER: 2006:267680 USPATFULL

TITLE: Dicer interacting proteins and uses therefor INVENTOR(S): Mello, Craig C., Shrewsbury, MA, UNITED STATES

PATENT ASSIGNEE(S): UNIVERSITY OF MASSACHUSETTS, Boston, MA, UNITED STATES (U.S. corporation)

NUMBER KIND DATE US 20060228361 A1 20061012 US 20070031417 A2 20070208 US 2005-107336 A1 20050414 (11) PATENT INFORMATION: APPLICATION INFO.:

> NUMBER DATE -----

PRIORITY INFORMATION: US 2004-562420P 20040414 (60) DOCUMENT TYPE: Utility

FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: LAHIVE & COCKFIELD, 28 STATE STREET, BOSTON, MA, 02109, US

NUMBER OF CLAIMS: 30 1

EXEMPLARY CLAIM: 11 Drawing Page(s) NUMBER OF DRAWINGS:

LINE COUNT: 11730 CAS INDEXING IS AVAILABLE FOR THIS PATENT.

Dicer (e.g., DCR-1) interactors are disclosed as are methods to positively or negatively modulate Dicer activity. Uses of Dicer interactors as drug targets are featured. Also featured are uses of Dicer interactors and modulators of same to modulate various Dicer

functions in vitro, in cell cultures, and in vivo.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 37 OF 160 CAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2006:1189497 CAPLUS

DOCUMENT NUMBER: 146:158125

TITLE: DUF283 domain of Dicer proteins has a double-stranded

RNA-binding fold Dlakic, Mensur

AUTHOR(S):

CORPORATE SOURCE: Department of Microbiology, Montana State University,

Bozeman, MT, 59717, USA SOURCE: Bioinformatics (2006), 22(22), 2711-2714

CODEN: BOINFP; ISSN: 1367-4803

PUBLISHER: Oxford University Press

DOCUMENT TYPE: Journal

LANGUAGE: English

Two RNases, Dicer and Argonaute, are at the heart of the RNA interference (RNAi) mol. machinery responsible for gene silencing. Both RNases contain multiple domains, most of which have been characterized or have functions that can be predicted based on sequence comparisons. However, Dicers of higher eukarvotes contain the domain known as DUF283 which at present has no assigned role. Using sensitive profile-profile comparisons, the authors detected a divergent double-stranded RNA-binding domain coinciding

with the DUF283 of Dicer. This finding has potential implications regarding the mechanistic role of Dicer in RNAi.

REFERENCE COUNT: THERE ARE 39 CITED REFERENCES AVAILABLE FOR THIS 39 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L6 ANSWER 95 OF 160 LIFESCI COPYRIGHT 2008 CSA on STN DUPLICATE 11

ACCESSION NUMBER: 2004:65552 LIFESCI

TITLE: Degradation of Double-Stranded RNA by Human Pancreatic

Ribonuclease: Crucial Role of Noncatalytic Basic Amino Acid Residues

AUTHOR: Sorrentino, S.; Naddeo, M.; Russo, A.; D'Alessio, G.

CORPORATE SOURCE: Department of Biological Chemistry, University Federico II of Naples, Naples, Italy

SOURCE: Biochemistry (Washington) [Biochemistry (Wash.)], (20030902

vol. 42, no. 34, pp. 10182-10190.

ISSN: 0006-2960.

DOCUMENT TYPE: Journal FILE SEGMENT:

LANGUAGE: English SUMMARY LANGUAGE: English

Under physiological salt conditions double-stranded

(ds) RNA is resistant to the action of most mammalian extracellular ribonucleases (RNases). However, some

pancreatic-type RNases are able to degrade

dsRNA under conditions in which the activity of boyine

RNase A, the prototype of the RNase superfamily, is

essentially undetectable. Human pancreatic ribonuclease (HP-

RNase) is the most powerful enzyme to degrade dsRNA within the tetrapod RNase superfamily, being

500-fold more active than the orthologous bovine enzyme on this substrate.

HP-RNase has basic amino acids at positions where RNase

A shows instead neutral residues. We found by modeling that some of these

basic charges are located on the periphery of the substrate binding site. To verify the role of these residues in the

cleavage of dsRNA, we prepared four variants of HP-

RNase: R4A, G38D, K102A, and the triple mutant R4A/G38D/K102A. The

overall structure and active site conformation of the variants were not

significantly affected by the amino acid substitutions, as deduced from CD

spectra and activity on single-stranded RNA

substrates. The kinetic parameters of the mutants with double

-helical poly(A) times poly(U) as a substrate were determined, as well as their helix-destabilizing action on a synthetic DNA substrate. The results

obtained indicate that the potent activity of HP-RNase on

dsRNA is related to the presence of noncatalytic basic residues

which cooperatively contribute to the binding and

destabilization of the double-helical RNA molecule.

These data and the wide distribution of the enzyme in different organs and body fluids suggest that HP-RNase has evolved to perform both

digestive and nondigestive physiological functions.

ANSWER 99 OF 160 USPATFULL on STN

ACCESSION NUMBER: 2002:294558 USPATFULL

TITLE: Human RNase III and compositions and uses thereof INVENTOR(S): Wu, Hongjiang, Carlsbad, CA, UNITED STATES

Crooke, Stanley T., Carlsbad, CA, UNITED STATES

NUMBER KIND DATE PATENT INFORMATION: US 20020164601 A1 20021107 US 6737512 B2 20040518 US 2001-900425 APPLICATION INFO .: A1 20010706 (9)

RELATED APPLN. INFO.: Continuation-in-part of Ser. No. US 2000-479783, filed

on 7 Jan 2000, PENDING Division of Ser. No. US 1997-870608, filed on 6 Jun 1997, PATENTED

Continuation-in-part of Ser. No. US 1996-659440, filed

on 6 Jun 1996, PATENTED DOCUMENT TYPE: Utility

FILE SEGMENT:

APPLICATION

LEGAL REPRESENTATIVE: John W. Caldwell, WOODCOCK WASHBURN LLP, One Liberty

Place-46th Floor, Philadelphia, PA, 19103

NUMBER OF CLAIMS: 52

EXEMPLARY CLAIM:

NUMBER OF DRAWINGS: 2 Drawing Page(s)

LINE COUNT: 1423

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The present invention provides polynucleotides encoding human RNAse III and polypeptides encoded thereby. Methods of using said polynucleotides

and polypeptides are also provided.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L6 ANSWER 113 OF 160 CAPLUS COPYRIGHT 2008 ACS on STN 2002:840867 CAPLUS

ACCESSION NUMBER: 138:119175

DOCUMENT NUMBER:

TITLE: Ribonuclease activity and RNA binding of recombinant

human Dicer

AUTHOR(S): Provost, Patrick; Dishart, David; Doucet, Johanne;

Frendewey, David; Samuelsson, Bengt; Radmark, Olof Department of Medical Biochemistry and Biophysics, CORPORATE SOURCE: Karolinska Institute, Stockholm, S-171 77, Swed. EMBO Journal (2002), 21(21), 5864-5874

SOURCE:

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RNA silencing phenomena, known as post-transcriptional gene silencing in plants, quelling in fungi, and RNA interference (RNAi) in animals, are mediated by double-stranded RNA (dsRNA) and mechanistically intersect at the RNase Dicer. Here, we report cloning and expression of the 218 kDa human Dicer, and characterization of its RNase

activity and dsRNA-binding properties. The recombinant enzyme generated .apprx.21-23 nucleotide products from dsRNA. Processing of the microRNA let-7 precursor by Dicer produced an apparently mature let-7 RNA. Mg2+ was required for dsRNase activity, but not for dsRNA binding, thereby uncoupling these reaction steps. ATP was dispensable for dsRNase activity in vitro. The Dicer-dsRNA complex formed at high KCl concns. was catalytically inactive, suggesting that lonic interactions are involved in dsRNA cleavage. The putative dsRNA-binding domain located at the C-terminus of Dicer was demonstrated to bind dsRNA in vitro. Human Dicer expressed in mammalian cells colocalized with calreticulin, a resident protein of the endoplasmic reticulum. A valiability of the recombinant Dicer protein will help improve our understanding of RNA silencing and other Dicer-related processes.

REFERENCE COUNT:

THERE ARE 52 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

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INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHBS, DIOTECHBS, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 11:59:20 ON 05 JUN 2008 SEA DICER? OR (DSRNAY(S)) (DEEGRAP2 OR CLEAVY)) OR ((DOUBLY(S) STRAN

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